```
SEQUENCE LISTING
         (1) GENERAL INFORMATION:
         (iii) NUMBER OF SEQUENCES: 5
         (2) INFORMATION FOR SEQ ID NO: 1:
5
         (i) SEQUENCE CHARACTERISTICS:
         (A) LENGTH: 323 amino acids
         (B) TYPE: amino acid
         (D) TOPOLOGY: linear
         (ii) MOLECULE TYPE: protein
10
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
         TyrSerArgG luLysAsnGlnProLysProSerProLysArgGluSer
         151015
         GlyGluGluPheArgMetGluLysLeuAsnGlnLeuTrpGluLysAla
         20 2530
15
         GlnArgLeuHisLeuProProValArgLeuAlaGluLeuHisAlaAsp
         354045
         LeuLysIleGlnGluArqAspGluLeuAl aTrpLysLysLeuLysLeu
         505560
         AspGlyLeuAspGluAspGlyGluLysGluAlaArgLeuIleArgAsn
20
         657075 80
         LeuAsnValIleLeuAlaLysTyrGlyLeuAspGlyLysLysAspAla
         859095
         ArgGlnValThrSerAsnSerLeuSerGlyThrGlnGluAspGly Leu
         100105110
25
         AspAspProArgLeuGluLysLeuTrpHisLysAlaLysThrSerGly
         115120125
         LysPheSerG lyGluGluLeuAspLysLeuTrpArgGluPheLeuHis
         130135140
         {\tt HisLysGluLysValHisGluTyrAsnValLeuLeuGluThrLeuSer}
30
         145150 155160
         ArgThrGluGluIleHisGluAsnValIleSerProSerAspLeuSer
         165170175
         AspIleLysGlySerValLeuHisSe rArgHisThrGluLeuLysGlu
         180185190
35
         {\tt LysLeuArgSerIleAsnGlnGlyLeuAspArgLeuArgArgValSer}
         195200 205
         {\tt HisGlnGlyTyrSerThrGluAlaGluPheGluGluProArgValIle}
         210215220
         {\tt AspLeuTrpAspLeuAlaGlnSerAlaAsnLeuThrAspLysGluLeu}
40
         225 230235240
         {\tt GluAlaPheArgGluGluLeuLysHisPheGluAlaLysIleGluLys}
         245250255
         HisAsnH isTyrGlnLysGlnLeuGluIleAlaHisGluLysLeuArg
         260265270
45
         {\tt HisAlaGluSerValGlyAspGlyGluArgValSerArgSerArgGlu}
         275 280285
         LysHisAlaLeuLeuGluGlyArgThrLysGluLeuGlyTyrThrVal
         290295300
         {\tt LysLysHisLeuGlnAspLeuSerGlyArgIleSe\ rArgAlaArgHis}
50
         305310315320
         AsnGluLeu
         (2) INFORMATION FOR SEQ ID NO: 2:
         (i) SEQUENCE CHARACTERISTICS:
         (A) LENGTH: 209 amino acids
55
         (B) TYPE: amino acid
         (D) TOPOLOGY: linear
         (ii) MOLECULE TYPE: protein
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
         ProArgLeuGluLysLeuTrpHisLysAlaLysThrSerGlyLysPhe
60
         151015
         SerGlyGluGluLeuAspLysLeuTrpArgGluPheLeu HisHisLys
```

```
202530
         GluLysValHisGluTyrAsnValLeuLeuGluThrLeuSerArgThr
         GluG luIleHisGluAsnValIleSerProSerAspLeuSerAspIle
 5
         LysGlySerValLeuHisSerArgHisThrGluLeuLysGluLysLeu
         657 07580
         ArgSerIleAsnGlnGlyLeuAspArgLeuArgArgValSerHisGln
10
         GlyTyrSerThrGluAlaGl uPheGluGluProArgValIleAspLeu
         100105110
         TrpAspLeuAlaGlnSerAlaAsnLeuThrAspLysGluLeuGluAla
         115120 125
         PheArgGluGluLeuLysHisPheGluAlaLysIleGluLysHisAsn
15
         130135140
         HisTyrGlnLysGlnLeuGluIleAlaHisGluLysLeuArgHisAla
         145150155160
         GluSerValGlyAspGlyGluArgValSerArgSerArgGluLysHis
         165170175
20
         A laLeuLeuGluGlyArgThrLysGluLeuGlyTyrThrValLysLys
         180185190
         HisLeuGlnAspLeuSerGlyArqIleSerArqAlaArqHisAsnGlu
         19 5200205
         Leu
25
         (2) INFORMATION FOR SEQ ID NO: 3:
         (i) SEQUENCE CHARACTERISTICS:
         (A) LENGTH: 33 base pairs
         (B) TYPE: nucleic acid
         (C) STRANDEDNESS: single
30
         (D) TOPOLOGY: linear
         (ii) MOLECULE TYPE: DNA (oligonucleotide)
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
         CCGCGTGGAT CCCCCAGGCTGGAAAAGCTGTGG33
         (2) INFORMATION FOR SEQ ID NO: 4:
35
         (i) SEQUENCE CHARACTERISTICS:
         (A) LENGTH: 35 base pairs
         (B) TYPE: nucleic acid
         (C) STRANDEDNESS: single
         (D) TOPOLOGY: linear
40
         (ii) MOLECULE TYPE: DNA (oligonucleotide)
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
         TCAATGAATTCTCAGAGTT CGTTGTGCCGAGCTCT35
         (2) INFORMATION FOR SEQ ID NO: 5:
         (i) SEQUENCE CHARACTERISTICS:
45
         (A) LENGTH: 205 amino acids
         (B) TYPE: amino acid
         (D) TOPOLOGY: linear
         (ii) MOLECULE TYPE: protein
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
50
         ProArgLeuGluLysLeuTrpHisLysAlaLysThr SerGlyIleSer
         151015
         ValArgLeuThrSerCysAlaArgValLeuHisTyrLysGluLysIle
         2025 30
         HisGluTyrAsnValLeuLeuAspThrLeuSerArgAlaGluGluGly
55
         TyrGluAsnLeuLeuSerProSerAspMetThrHisIleLysSerAsp
         50 5560
         ThrLeuAlaSerLysHisSerGluLeuLysAspArgLeuArgSerIle
         65707580
60
         AsnGlnGlyLeuAspArg LeuArgLysValSerHisGlnLeuArgPro
         859095
```

 ${\tt AlaThrGluPheGluGluProArgValIleAspLeuTrpAspLeuAla}$ 100 105110 ${\tt GlnSerAlaAsnPheThrGluLysGluLeuGluSerPheArgGluGlu}$ 115120125 5 LeuLysHisPheGluAlaLysIleGluLysHisAsn HisTyrGlnLys 130135140 GlnLeuGluIleSerHisGlnLysLeuLysHisValGluSerIleGly 145150155160 ${\tt AspProGluHisIleSerArgAsnLysGluLysTyrValLeuLeuGlu}$ 10 165170175 ${\tt GluLysThrLysGluLeuGlyTyrLysValLysLysHisLeuGlnAsp}$ 180185190 LeuSerSerArgValSerArgAlaArgHisAsnGluLeu 195200

The above description is meant to be illustrative of the present invention, and not limiting thereof. All explanations of the inventors theory of the invention are for illustrative purposes only. It is the inventors' intention that the scope of their invention be defined solely by the following claims.